

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.**

Application Serial Number: 10/507,140A  
Source: PCT  
Date Processed by STIC: 3/9/06

# ***ENTERED***



PCT

## RAW SEQUENCE LISTING

DATE: 03/09/2006

PATENT APPLICATION: US/10/507,140A

TIME: 12:23:16

Input Set : E:\PUS\_E005\_105b.txt

Output Set: N:\CRF4\03092006\J507140A.raw

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3 <110> APPLICANT: NERI, Dario
4     MELKKO, Samu
6 <120> TITLE OF INVENTION: Encoded Self-Assembling Chemical Libraries
(Esachel)
8 <130> FILE REFERENCE: PUS-E005-105B
10 <140> CURRENT APPLICATION NUMBER: 10/507,140A
C--> 11 <141> CURRENT FILING DATE: 2004-09-07
13 <150> PRIOR APPLICATION NUMBER: PCT/EP02/04153
14 <151> PRIOR FILING DATE: 2002-03-08
16 <160> NUMBER OF SEQ ID NOS: 28
18 <170> SOFTWARE: PatentIn version 3.3
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 60
22 <212> TYPE: DNA
23 <213> ORGANISM: Artificial Sequence
25 <220> FEATURE:
26 <223> OTHER INFORMATION: Primer L19VH_Eco_fo
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32 <210> SEQ ID NO: 2
33 <211> LENGTH: 66
34 <212> TYPE: DNA
35 <213> ORGANISM: Artificial Sequence
37 <220> FEATURE:
38 <223> OTHER INFORMATION: Primer L19VH_Hind_ba
40 <400> SEQUENCE: 2
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43 cagggt      66
46 <210> SEQ ID NO: 3
47 <211> LENGTH: 63
48 <212> TYPE: DNA
49 <213> ORGANISM: Artificial Sequence
51 <220> FEATURE:
52 <223> OTHER INFORMATION: Primer L19VL_Eco_fo
54 <400> SEQUENCE: 3
55 ttccacacag aattcattaa agaggagaaa ttaactatgg aaattgtgtt gacgcagtct      60
57 cca      63
60 <210> SEQ ID NO: 4
61 <211> LENGTH: 69
62 <212> TYPE: DNA
63 <213> ORGANISM: Artificial Sequence
65 <220> FEATURE:
66 <223> OTHER INFORMATION: Primer L19VL_Hind_ba
68 <400> SEQUENCE: 4

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69 tcaatctgat taagcttagt gatggtgatg gtgatgacat ccacctttga tttccacctt      60
71 ggtcccttg      69
74 <210> SEQ ID NO: 5
75 <211> LENGTH: 60
76 <212> TYPE: DNA
77 <213> ORGANISM: Artificial Sequence
79 <220> FEATURE:
80 <223> OTHER INFORMATION: Primer HH10VH_Eco_fo
82 <400> SEQUENCE: 5
83 tttcacacag aattcattaa agaggagaaa ttaactatgg aggtgaagct gcagcagtct      60
86 <210> SEQ ID NO: 6
87 <211> LENGTH: 66
88 <212> TYPE: DNA
89 <213> ORGANISM: Artificial Sequence
91 <220> FEATURE:
92 <223> OTHER INFORMATION: Primer HH10VH_Hind_ba
94 <400> SEQUENCE: 6
95 tcaatctgat taagcttagt gatggtgatg gtgatgacat ccacctgcag agacagtgc      60
97 cagagt      66
100 <210> SEQ ID NO: 7
101 <211> LENGTH: 63
102 <212> TYPE: DNA
103 <213> ORGANISM: Artificial Sequence
105 <220> FEATURE:
106 <223> OTHER INFORMATION: Primer HH10VL_Eco_fo
108 <400> SEQUENCE: 7
109 tttcacacag aattcattaa agaggagaaa ttaactatgg atattgtgct aactcagtct      60
111 cca      63
114 <210> SEQ ID NO: 8
115 <211> LENGTH: 69
116 <212> TYPE: DNA
117 <213> ORGANISM: Artificial Sequence
119 <220> FEATURE:
120 <223> OTHER INFORMATION: Primer HH10VL_Hind_ba
122 <400> SEQUENCE: 8
123 tcaatctgat taagcttagt gatggtgatg gtgatgacat ccacctttta tttccagctt      60
125 ggtccccc      69
128 <210> SEQ ID NO: 9
129 <211> LENGTH: 48
130 <212> TYPE: DNA
131 <213> ORGANISM: Artificial Sequence
133 <220> FEATURE:
134 <223> OTHER INFORMATION: with 5' -thiol
136 <400> SEQUENCE: 9
137 ggagcttctg aattctgtgt gctgcataat cgacacgaat tccgcagc      48
140 <210> SEQ ID NO: 10
141 <211> LENGTH: 48
142 <212> TYPE: DNA
143 <213> ORGANISM: Artificial Sequence

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145 <220> FEATURE:
146 <223> OTHER INFORMATION: with 3' -thiol
148 <400> SEQUENCE: 10
149 tcgcgagggg aattcgtcat atatcagcac acagaattca gaagctcc 48
152 <210> SEQ ID NO: 11
153 <211> LENGTH: 48
154 <212> TYPE: DNA
155 <213> ORGANISM: Artificial Sequence
157 <220> FEATURE:
158 <223> OTHER INFORMATION: HyHel10_5SH with 5'-thiol
160 <400> SEQUENCE: 11
161 ggagcttctg aattctgtgt gctgcagtgg cgacacgaat tccgcagc 48
164 <210> SEQ ID NO: 12
165 <211> LENGTH: 48
166 <212> TYPE: DNA
167 <213> ORGANISM: Artificial Sequence
169 <220> FEATURE:
170 <223> OTHER INFORMATION: Primer HyHel10_3SH with 3'-thiol
172 <400> SEQUENCE: 12
173 tcgcgagggg aattcgtcat agggcagcac acagaattca gaagctcc 48
176 <210> SEQ ID NO: 13
177 <211> LENGTH: 48
178 <212> TYPE: DNA
179 <213> ORGANISM: Artificial Sequence
181 <220> FEATURE:
182 <223> OTHER INFORMATION: Primer GST_5SH with 5'-thiol
184 <400> SEQUENCE: 13
185 ggagcttctg aattctgtgt gctgctgagg cgacacgaat tccgcagc 48
188 <210> SEQ ID NO: 14
189 <211> LENGTH: 48
190 <212> TYPE: DNA
191 <213> ORGANISM: Artificial Sequence
193 <220> FEATURE:
194 <223> OTHER INFORMATION: Primer GST_3SH with 3'-thiol
196 <400> SEQUENCE: 14
197 tcgcgagggg aattcgtcaa gaggcagcac acagaattca gaagctcc 48
200 <210> SEQ ID NO: 15
201 <211> LENGTH: 24
202 <212> TYPE: DNA
203 <213> ORGANISM: Artificial Sequence
205 <220> FEATURE:
206 <223> OTHER INFORMATION: Primer 1AB_PCRfo
208 <400> SEQUENCE: 15
209 ggagcttctg aattctgtgt gctg 24
212 <210> SEQ ID NO: 16
213 <211> LENGTH: 18
214 <212> TYPE: DNA
215 <213> ORGANISM: Artificial Sequence
217 <220> FEATURE:

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218 <223> OTHER INFORMATION: Primer 1A_PCRba
220 <400> SEQUENCE: 16
221 gctgcggaat tcgtgtcg                                     18
224 <210> SEQ ID NO: 17
225 <211> LENGTH: 18
226 <212> TYPE: DNA
227 <213> ORGANISM: Artificial Sequence
229 <220> FEATURE:
230 <223> OTHER INFORMATION: Primer 1B_PCRba
232 <400> SEQUENCE: 17
233 tcgcgagggg aattcgtc                                     18
236 <210> SEQ ID NO: 18
237 <211> LENGTH: 39
238 <212> TYPE: DNA
239 <213> ORGANISM: Artificial Sequence
241 <220> FEATURE:
242 <223> OTHER INFORMATION: Primer typeB_oligo
244 <400> SEQUENCE: 18
245 gcataccgga attcccagca taatgatcgc tatcgctgc           39
248 <210> SEQ ID NO: 19
249 <211> LENGTH: 63
250 <212> TYPE: DNA
251 <213> ORGANISM: Artificial Sequence
253 <220> FEATURE:
254 <221> NAME/KEY: modified_base
255 <222> LOCATION: (40)..(45)
256 <223> OTHER INFORMATION: Primer typeA_oligo with spacer elements n; n =
deoxyribose backbone elements without bases
258 <400> SEQUENCE: 19
259 cgtcagctcg aattctccat atatgcagcg atagcgatcd ddddctggg aattccggta   60
261 tgc                                                         63
264 <210> SEQ ID NO: 20
265 <211> LENGTH: 18
266 <212> TYPE: DNA
267 <213> ORGANISM: Artificial Sequence
269 <220> FEATURE:
270 <223> OTHER INFORMATION: Primer CodeABfo
272 <400> SEQUENCE: 20
273 gcataccgga attcccag                                     18
276 <210> SEQ ID NO: 21
277 <211> LENGTH: 18
278 <212> TYPE: DNA
279 <213> ORGANISM: Artificial Sequence
281 <220> FEATURE:
282 <223> OTHER INFORMATION: Primer CodeABba
284 <400> SEQUENCE: 21
285 cgtcagctcg aattctcc                                     18
288 <210> SEQ ID NO: 22
289 <211> LENGTH: 29
290 <212> TYPE: DNA

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Input Set : E:\PUS\_E005\_105b.txt

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291 <213> ORGANISM: Artificial Sequence

293 <220> FEATURE:

294 <223> OTHER INFORMATION: Primer with 5' sequence specifically coding a member of a

295       sublibrary; where n is a, c, g, or t

298 <220> FEATURE:

299 <221> NAME/KEY: misc\_feature

300 <222> LOCATION: (1)..(5)

301 <223> OTHER INFORMATION: n is a, c, g, or t

303 <400> SEQUENCE: 22

W--> 304 nnnnncagca cacagaattc agaagctcc 29

307 <210> SEQ ID NO: 23

308 <211> LENGTH: 29

309 <212> TYPE: DNA

310 <213> ORGANISM: Artificial Sequence

312 <220> FEATURE:

313 <223> OTHER INFORMATION: Primer with a 3' sequence specific for a member of a library;

314       where n is a, c, g, or t

317 <220> FEATURE:

318 <221> NAME/KEY: misc\_feature

319 <222> LOCATION: (25)..(29)'

320 <223> OTHER INFORMATION: n is a, c, g, or t

322 <400> SEQUENCE: 23

W--> 323 ggagcttctg aattctgtgt gctgnnnnn 29

326 <210> SEQ ID NO: 24

327 <211> LENGTH: 24

328 <212> TYPE: DNA

329 <213> ORGANISM: Artificial Sequence

331 <220> FEATURE:

332 <223> OTHER INFORMATION: Primer with 5' sequence specific for a chemical moiety linked to

333       primer by a biotinylated base analog

335 <400> SEQUENCE: 24

336 cagcacacag aattcagaag ctcc 24

339 <210> SEQ ID NO: 25

340 <211> LENGTH: 48

341 <212> TYPE: DNA

342 <213> ORGANISM: Artificial Sequence

344 <220> FEATURE:

345 <223> OTHER INFORMATION: Primer with 5' iminobiotinyl-NH-(CH<sub>2</sub>)<sub>6</sub> group

347 <400> SEQUENCE: 25

348 ggagcttctg aattctgtgt gctgattggc cgacacgaat tccgcagc 48

351 <210> SEQ ID NO: 26

352 <211> LENGTH: 48

353 <212> TYPE: DNA

354 <213> ORGANISM: Artificial Sequence

356 <220> FEATURE:

357 <223> OTHER INFORMATION: Primer with 3' (CH<sub>2</sub>)<sub>6</sub>-NH-iminobiotinyl group

359 <400> SEQUENCE: 26

360 tcgcgagggg aattcgtcat ttaccagcac acagaattca gaagctcc 48

363 <210> SEQ ID NO: 27

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/10/507,140A

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TIME: 12:23:17

Input Set : E:\PUS\_E005\_105b.txt  
Output Set: N:\CRF4\03092006\J507140A.raw

**Please Note:**

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220>

to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:22; N Pos. 1,2,3,4,5 ✓

Seq#:23; N Pos. 25,26,27,28,29 ✓

**Invalid Line Length:**

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:19; Line(s) 256

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/10/507,140A

DATE: 03/09/2006

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Input Set : E:\PUS\_E005\_105b.txt

Output Set: N:\CRF4\03092006\J507140A.raw

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:304 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0

L:323 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0